

PubMed Nucleotide Protein Genome Structure PopSet Taxonomy OMIM Bio

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☐ 1: AW512144. xu54e11.x1 NCI_CG...[gi:7150222]

[NEW](#) [Links](#)

LOCUS AW512144 468 bp mRNA linear EST 03-MAR-2000
 DEFINITION xu54e11.x1 NCI_CGAP Ut1 Homo sapiens cDNA clone IMAGE:2805548 3' similar to SW:ES10_RAT P16303 LIVER CARBOXYLESTERASE 10 PRECURSOR ;, mRNA sequence.
 ACCESSION AW512144
 VERSION AW512144.1 GI:7150222
 KEYWORDS EST.
 SOURCE human.
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo/Pan/Gorilla group; Homo.
 REFERENCE 1 (bases 1 to 468)
 AUTHORS NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>.
 TITLE National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index
 JOURNAL Unpublished (1997)
 COMMENT Contact: Robert Strausberg, Ph.D.
 Email: cgapbs-r@mail.nih.gov
 Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D.
 cDNA Library Preparation: Life Technologies, Inc.
 cDNA Library Arrayed by: Greg Lennon, Ph.D.
 DNA Sequencing by: Washington University Genome Sequencing Center
 Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov/image/html/iresources.shtml

Possible reversed clone: similarity on wrong strand

Seq primer: -40UP from Gibco




High quality sequence stop: 406.

FEATURES Location/Qualifiers
 source 1..468
 /organism="Homo sapiens"
 /db_xref="taxon:9606"
 /clone="IMAGE:2805548"
 /clone_lib="NCI_CGAP_Ut1"
 /tissue_type="well-differentiated endometrial adenocarcinoma, 7 pooled tumors"
 /lab_host="DH10B"
 /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.75 kb. Life Technologies catalog #: 11538-014"

BASE COUNT 72 a 141 c 162 g 89 t 4 others

ORIGIN

1 ctgcaggagt cctggggcca gctggcctcg atgtacgtca gcacgcggga acggtacaag
 61 tggctgcgct tcagcgagga ctgtctgtac ctgaacgtgt acgcgcgggc ggcgcgcgcc

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1 ctgcaggagt cctggggcca gctggcctcg atgtacgtca gcacgcggga acggtacaag
 61 tggctgcgct tcagcgagga ctgtctgtac ctgaacgtgt acgcgccggc gcgcgcgcc

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241 cagcacaggc tcggcatctt cggcttcctg agcacggacg acagccacgc gcgcgggaac
301 tgggggctgc tggaccagat ggcggctctg cgctgggtgc aggagaacat cgcagccttn
361 cgggggagacc cangaaatgt gacctgttc ggccagtcgg cggggggccat gagcatctca
421 ggactgatga tgnacccta gctcgggct cttcatcngg ccatttcc
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Revised: July 5, 2002.

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Aug 28 2002 15:52:55